

FIGURE 1

|     |  |     |
|-----|--|-----|
| 1   | ACTGTGGACTTTCTCTGGGGTGTCTGGTCAATGCCCTTACAGTATGGTGAGATCTGCTGAG        | 60  |
| 61  | CACCTGTGGTAATTTGGAGAAAGTCTCTCTGTAATAATTACACACAAGCACCGACATTAATGCTG    | 120 |
| 121 | AGCTCAGCCCTCCAAATTTCCAAATTTGTCCTTTCATCTCCAAATTCACCCGCTACTATGCTGTGTGT | 180 |
| 181 | GATCCACTGAGATAAAGCCCAAGATGAATATCTTTGGTTAATTTGTGTGATGATCTTTCATT       | 240 |
| 241 | AGTTGGAGTGTCCCTGCTGTCTTTTGCAATTTGGAAATGATCTTTCTGGAGCTAAACCTCAAA      | 300 |
| 301 | GGCGCTGAAGAGATAATACAAACATGTTCACATGCAGAGGAGGTTGCTCTGTCTCTCTTT         | 360 |
| 361 | AGCAAAATATCTGGGGTACTGACCTTTATGACTTCTTTTATATACCTGGATCTATTATG          | 420 |
| 421 | TTATGTGCTATTTACAGAAATAATCTTTATTCGCTAAAGAACAGCAAGATTAAATTAGTGAT       | 480 |
| 481 | GCCAAATCAGAAGCTCCAAATTTGGAAATTCGAAATTCGAAATTTTCACAAAGCAAGAA          | 540 |
| 541 | AGGAAAGCTTGTGAAGACAAATTTGGGGATTTGTGATG                               | 573 |

[illegible]

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FIGURE 3A

1   ATTGCTCGACAGCCAAGGACAGAGCAGCCCTGTGTTTAGTTCTCTGTAGTATGCATCT   60  
61   TTGCCACAATAGCGCGAATATTTCCCAACAGCAAC'GGTCAAGGATGTCCGTGC   120  
121   TTCGCTGTACAGCTTAATATCACT'CA'ATAATTCTAACCACTCTGGTTGGCAACTTAATAGT   180  
181   AATCA'TTTCGATATCCCCACT'FCAAGCAACTTCACACGCCCAACAATTTGGCTCCTTCATTTC   240  
241   CATGCCCGTTGTCGACTTTTCTGTCTGGGCTGTCTGGTCAATGCCCTACAGCATGGTGAGAAC   300  
301   AGTTGAGCACTGCTGGTACTTTTGGGGAAC'TCTCTGCAAACTTCACACCAGCACTGATAT   360  
361   CATGCTGAGCTCGGCATCCA'TTCTCCACCTAGCC'TTCA'TTCCATTTGACCCGCTACTATGC   420  
421   TGTGTGCGACCC'TTAAAGATACAAAGCCAAGATCAATCTCGCCGCCATTTTGTGTGATGAT   480  
481   CCTCATTAGCTGGAGCCCTTCCCTGCTGTTT'TTGCA'TTTGGGATGATCTTCCCTGGAGCTGAA   540  
541   CTTAGAAGGAGTTGAGGAGCTGTATCACAAATCAGGTCTCTGCCCTGCGCGGCTGTTTTC   600  
601   CTTCTTCAGTAAAGTATCTGGGGTACTGGCATTCATGACGTCTTCTATATATACCTGGATC   660

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## FIGURE 3B

|      |  |      |
|------|--|------|
| 661  | TGTTATGTTATTGTTTACTATAGAAATATATTCATAGCTAAAGGACAAGCAAGGTCAAT    | 720  |
| 721  | TAATCGTGCAAAATCTTCAAGTTGGATTTGGAAGGGAAAGCAGAGCGCCACAAAGCAAGGA  | 780  |
| 781  | AACAAAGCCGCGAAACCTTAGGGATCATGGTGGCGTTTTCCTCCTGTGCTGGTGCCCC     | 840  |
| 841  | GTTCCTTTTCTGCA TGTCCTGGACCCCTTTCCTGGGCTATGTTATCCCACTCTCTGAA    | 900  |
| 901  | TGACACACTGAAATGGTTTGGGTACCTGAACTCTGCCCTTCAACCCGATGGTTTATGCCCTT | 960  |
| 961  | TTTCTATCCCTGGTTCAGAAAGAGCGTTGAAGATGGTTCCTTCGGTAAATAATTTTCCAAAA | 1020 |
| 1021 | AGATTCATCTAGGTCCTAAGTTATTTTGTAAACGCAATCCATGAACCAAGTATATTTTGTA  | 1080 |
| 1081 | GTTCCTTAAGAGCAGTTGGTGA   | 1101 |

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## FIGURE 4A

|     |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |     |     |
|-----|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|-----|-----|
| 1   | M | H | L | C | H | N | S | A | N | I | S | H | T | N | S | N | W | S | R | D   | 20  |
| 21  | V | R | A | S | L | Y | S | L | I | S | L | I | I | L | T | T | L | V | G | N   | 40  |
| 41  | L | I | V | I | I | S | I | S | H | F | K | Q | L | H | T | P | T | N | W | L   | 60  |
| 61  | L | H | S | M | A | V | V | D | F | L | L | G | C | L | V | M | P | Y | S | M   | 80  |
| 81  | V | R | T | V | E | H | C | W | Y | F | G | E | L | F | C | K | L | H | T | S   | 100 |
| 101 | T | D | I | M | L | S | S | A | S | I | L | H | L | A | F | I | S | I | D | R   | 120 |
| 121 | Y | Y | A | V | C | D | P | L | R | Y | K | A | K | I | N | L | A | A | I | F   | 140 |
| 141 | V | M | I | L | I | S | W | S | L | P | A | V | F | A | F | G | M | I | F | L   | 160 |
| 161 | E | L | N | L | E | G | V | E | E | L | Y | H | N | Q | V | F | C | L | R | G   | 180 |
| 181 | C | F | P | F | S | K | V | S | G | V | L | A | F | M | T | S | F | Y | I | 200 |     |
| 201 | P | G | S | V | M | L | F | V | Y | Y | R | I | Y | F | I | A | K | G | Q | A   | 220 |

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## FIGURE 4B

221 R S I N R A N L Q V G L E G E S R A P Q 240  
241 S K E T K A A K T L G I M V G V F L L C 260  
261 W C P F F C M V L D P F L G Y V I P F 280  
281 T L N D T L N W F G Y L N S A F N P M V 300  
301 Y A F F Y P W F R R A L K M V L F G K I 320  
321 F Q K D S S R S K L F L 332

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FIGURE 5A

1 TCAGGAATGATGCCCTTTTGCCACAATATAATTAATATTTCCCTGTGTGAAAAACAACACTGG 60  
61 TCAAAATGATGTCCTGCTTCCCTGTACAGTTTAAATGGTGCTCATAATTCTGACCACACTC 120  
121 GTTGGCAATCTGATAGTTATTGTTTCTATATCACACTTCAAACAACACTTCATACCCCAACA 180  
181 AATGGCTCATTCATCCATGGCCACTGTGGACTTTCCTTCTGGGGTGCTGGTCAATGCCT 240  
241 TACAGTATGGTGAGATCTGCTGAGCACTGTTGGTATTTTGGAGAAGTCTTCTGTAAAATT 300  
301 CACACAAGCACCGACATTATGCTGAGCTCAGCCCTCCATTTCCTCATTTTGTCTTTCATCTCC 360  
361 ATTGACCGCTACTATGCTGTGTGATCCACTGACATATAAAGCCAAAGATGAATATCTTG 420  
421 GTTATTTGTGATGATCTTCATTAGTTGGAGTGTCCTGCTGTTTTCGATTTTGGAAATG 480  
481 ATCTTTCTGGAGCTAAACTTCAAAGGCGCTGAAGAGATATATTACAAACATGTTCACTGC 540  
541 AGAGGAGGTTGGCTCTGCTCTTCTTTAGCAAAATATCTGGGGTACTGACCTTTATGACTTCT 600  
601 TTTTATATACCTGGATCTATTATGTTATGTGCTATTACAGAATATATCTTATCGCTAAA 660

## FIGURE 5B

661 GAACAGGCAAGATTAATTAGTGATGCCAATCAGAAGCTCCAATTTGGATTGGAAATGAAA 720  
721 AATGGAATTCACAAAGCAAGAAAGAAAGCTGTGAAGACATTGGGGATTGTGATGGGA 780  
781 GTTTTCCTAATATGCTGGTGCCCTTTCTTTATCTGTACAGTCATGGACCCCTTTTCTTCAC 840  
841 TACATTATCCACCCTACTTTGAATGATGTGTTGATTGGTTGGCTACTTGAACTCTACA 900  
901 TTTAATCCAATGGTTTATGCATTTTCTATCCTTGGTTTAGAAAAGCACTGAAGATGATG 960  
961 CTGTTGGTAAAATTTCCAAAAGATTATCCAGGTGTAATATTTTGGAAATGAGT 1020  
1021 TCATAGAATTATATATT 1038



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FIGURE 6A

|     |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |     |
|-----|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|-----|
| 1   | M | M | P | F | C | H | N | I | I | N | I | S | C | V | K | N | N | W | S | N | 20  |
| 21  | D | V | R | A | S | L | Y | S | L | M | V | L | I | I | L | T | T | L | V | G | 40  |
| 41  | N | L | I | V | I | V | S | I | S | H | F | K | Q | L | H | T | P | T | N | W | 60  |
| 61  | L | I | H | S | M | A | T | V | D | F | L | L | G | C | L | V | M | P | Y | S | 80  |
| 81  | M | V | R | S | A | E | H | C | W | Y | F | G | E | V | F | C | K | I | H | T | 100 |
| 101 | S | T | D | I | M | L | S | S | A | S | I | F | H | L | S | F | I | S | I | D | 120 |
| 121 | R | Y | Y | A | V | C | D | P | L | R | Y | K | A | K | M | N | I | L | V | I | 140 |
| 141 | C | V | M | I | F | I | S | W | S | V | P | A | V | F | A | F | G | M | I | F | 160 |
| 161 | L | F | L | N | F | K | G | A | E | E | I | Y | Y | K | H | V | H | C | R | G | 180 |
| 181 | G | C | S | V | F | F | S | K | I | S | G | V | L | T | F | M | T | S | F | Y | 200 |
| 201 | I | P | G | S | I | M | L | C | V | Y | Y | R | I | Y | L | I | A | K | E | Q | 220 |

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FIGURE 6B

221 A R L I S D A N Q K L Q I G L E M K N G 240  
241 I S Q S K E R K A V K T L G I V M G V F 260  
261 L I C W C P F F I C T V M D P F L H Y I 280  
281 I P P T L N D V L I W F G Y L N S T F N 300  
301 P M V Y A F F Y P W F R K A L K M M L F 320  
321 G K I F Q K D S S R C K L F L E L S S \* 340

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FIGURE 7A

|             |     |   |     |
|-------------|-----|---|-----|
| Rat SNORF33 | 1   | .MHLCHNSANISHTNSNWSRDVRASLYSLISLIILTTLVGNLIVISISH   | 49  |
|             |     | .           .           .           .               |     |
| Hum SNORF33 | 1   | MMPECHNIINISCVKNNWSNDVRASLYSLMVLIILTTLVGNLIVISISH   | 50  |
| Rat SNORF33 | 50  | FKQLHTPTNWLHSMVVDVFLGCLVMPYSMVRTVEHCWYFGEFCKLHT     | 99  |
|             |     | :           .           .           .               |     |
| Hum SNORF33 | 51  | FKQLHTPTNWLHSMATVDVFLGCLVMPYSMVRSAEHCWYFGEVFCKIHT   | 100 |
| Rat SNORF33 | 100 | STDIMLSSASILHLAFISIDRYYAVCDPLRYKAKINLAAIFVMILISWSL  | 149 |
|             |     | :           .                                       |     |
| Hum SNORF33 | 101 | STDIMLSSA:IFHLFSIDRYYAVCDPLRYKAKMNILVICVMIFISWSV    | 150 |
| Rat SNORF33 | 150 | PAVFAFGMIFLELNLEGVVEELYHNQVFCRLRGCFPFSSKVSGLAFMTSFY | 199 |
|             |     |   |     |
| Hum SNORF33 | 151 | PAVFAFGMIFLELNFKGAEIYKHHVHCRGCGSVFFSKISGVLTFMTSFY   | 200 |
| Rat SNORF33 | 200 | IPGSVMLFVYYRIYFIAGQARSINRAN..LQVGLEGESRAPQSKETKAA   | 247 |
|             |     | .   |     |
| Hum SNORF33 | 201 | IPGSIMLCVYYRIYLIAKEQARLISDANQKLQIGLEMKNGISQSKERKAV  | 250 |

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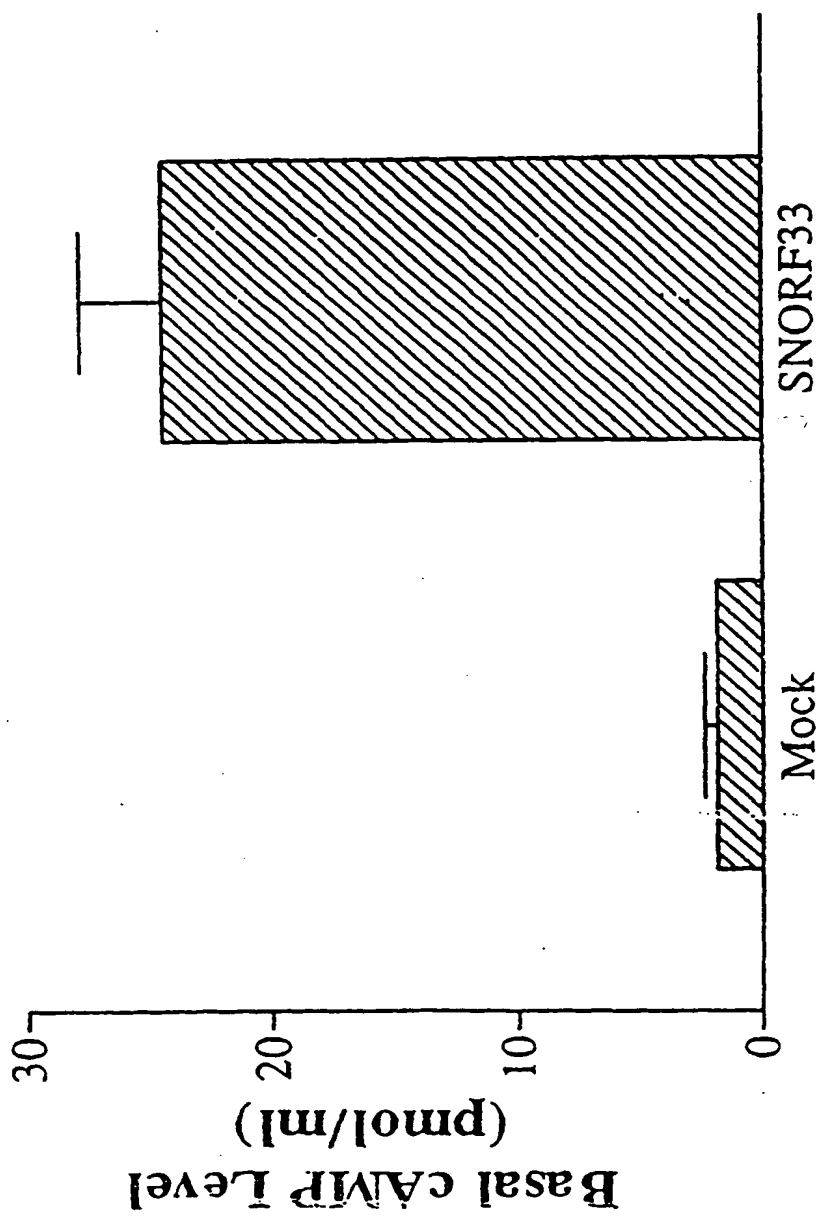
## FIGURE 7B

|             |     |   |     |
|-------------|-----|---|-----|
| Rat SNORF33 | 248 | KTGIMVGVFLLCWCPEFFFCMVLDPELGYVIPPTLNDTLNWFYLN | 297 |
|             |     | .    :          :          :                  |     |
| Hum SNORF33 | 251 | KTGIVMGVFLICWCPEFFICTVMDPELHYIIPPTLNDVLIWFYLN | 300 |
|             |     |   |     |
| Rat SNORF33 | 298 | PMVYAFFYPWFRRALKMVLEFGKIFQKDSRSRKLFL....*     | 333 |
|             |     | :     .     :                                 |     |
| Hum SNORF33 | 301 | PMVYAFFYPWFRKALKMMLFGKIFQKDSRSRCKLFLELSS*     | 340 |

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FIGURE 8

Basal cAMP Levels in Mock- and rat  
SNORF33-Transfected Cos-7 cells



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FIGURE 9

Agonist-Mediated Increase in Intracellular cAMP Levels in  
Mock- and rat SNORF33-Transfected Cos-7 Cells

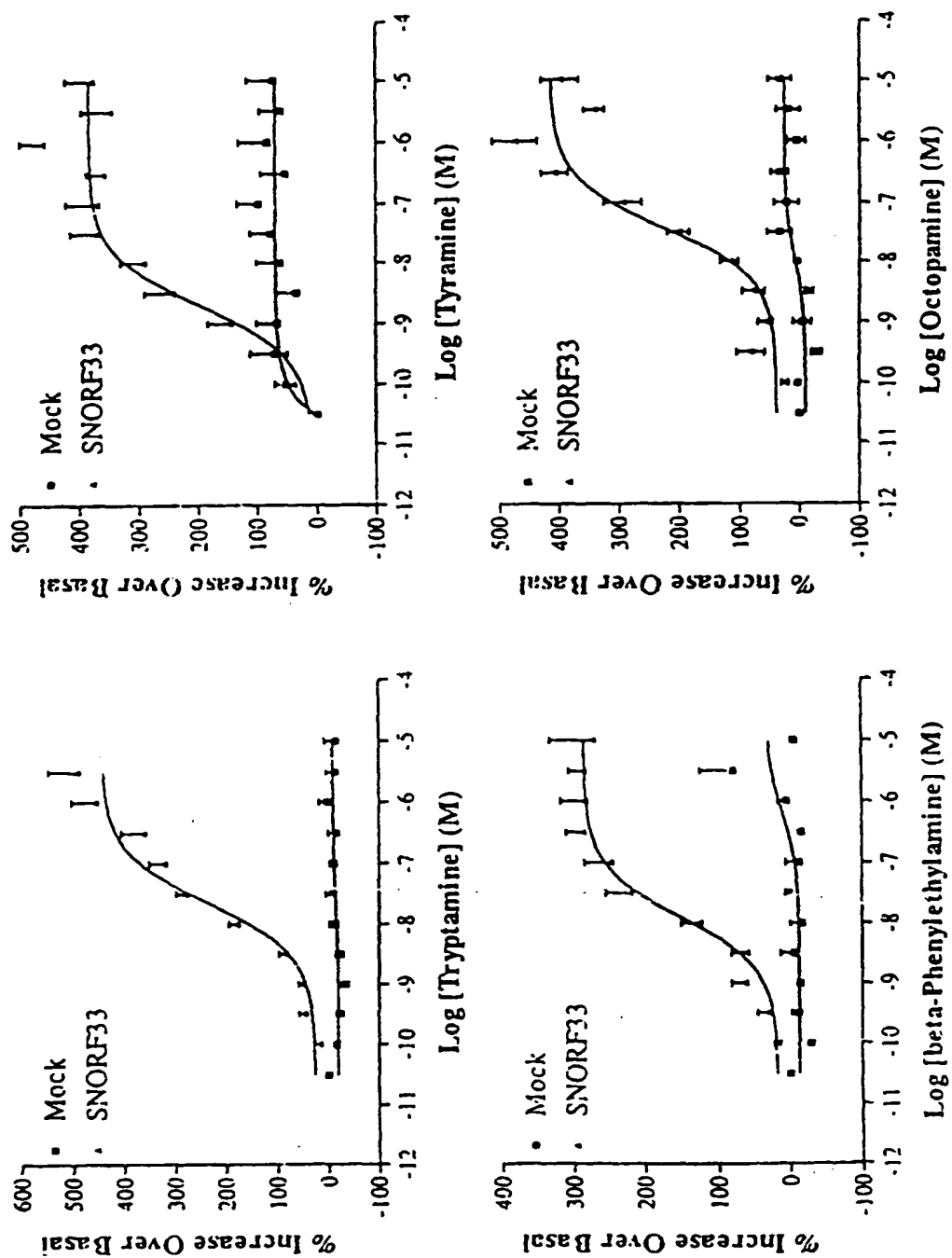
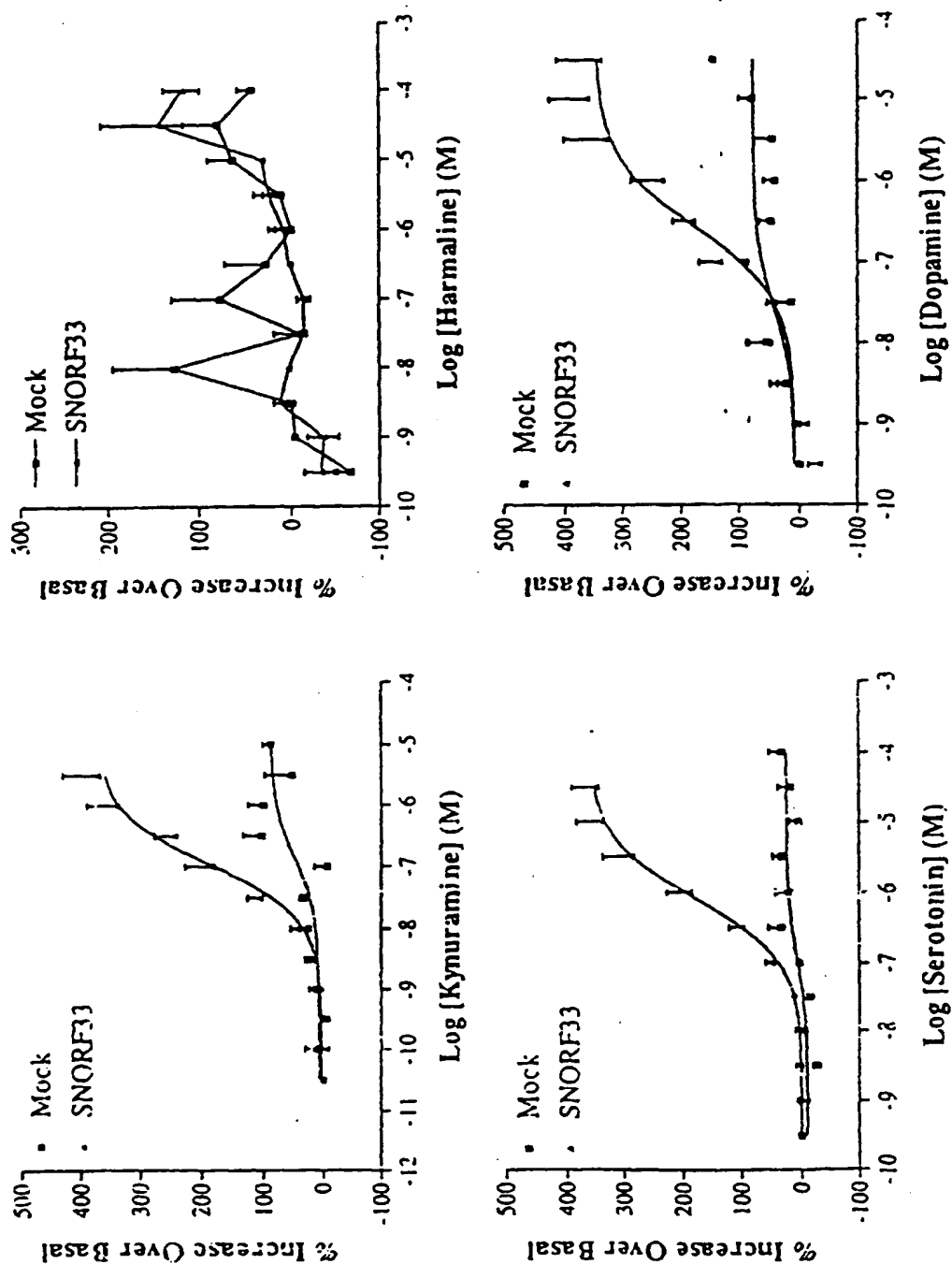


FIGURE 10  
Agonist-Mediated Increase in Intracellular cAMP Levels in  
Mock- and rat SNORF33-Transfected Cos-7 Cells



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Figure 11A

Octopamine

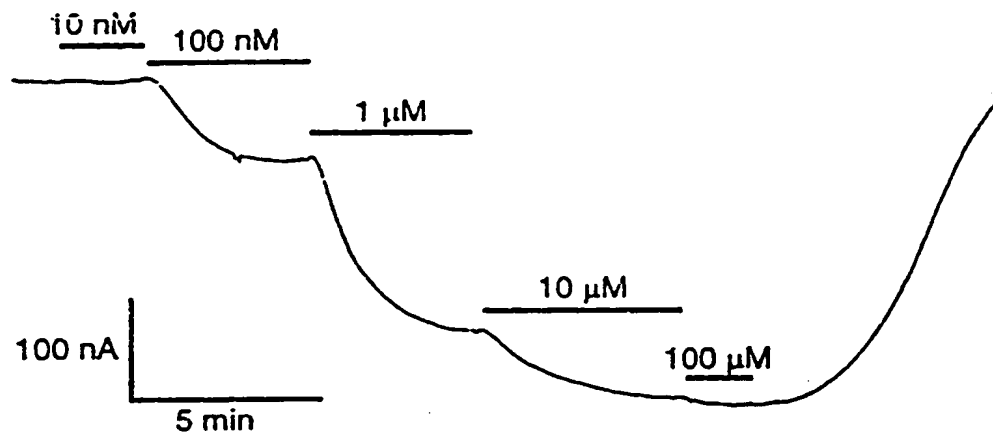
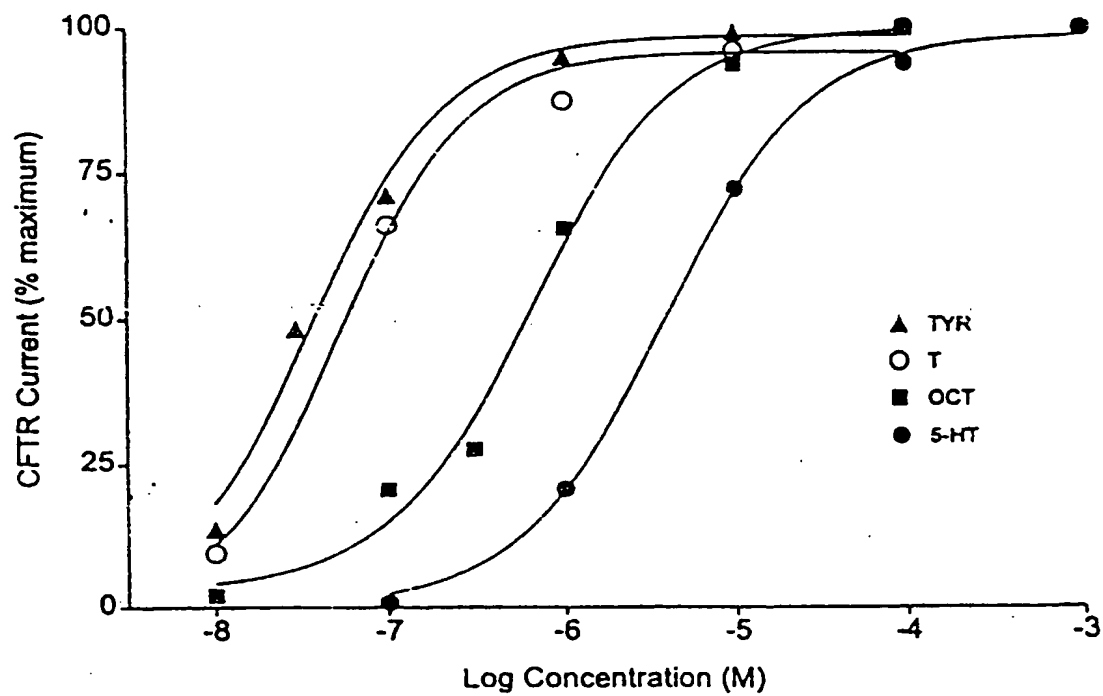
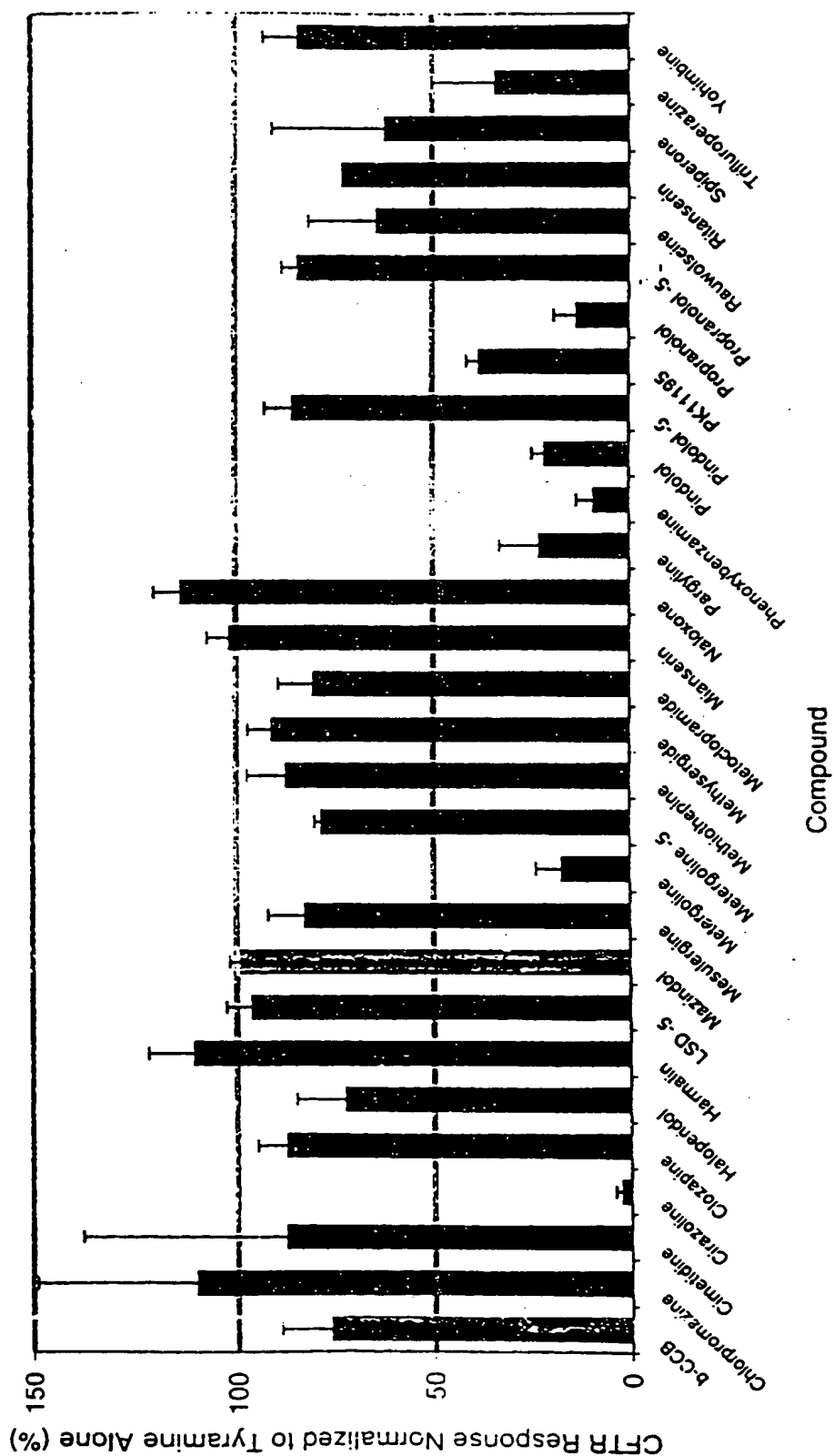


Figure 11B

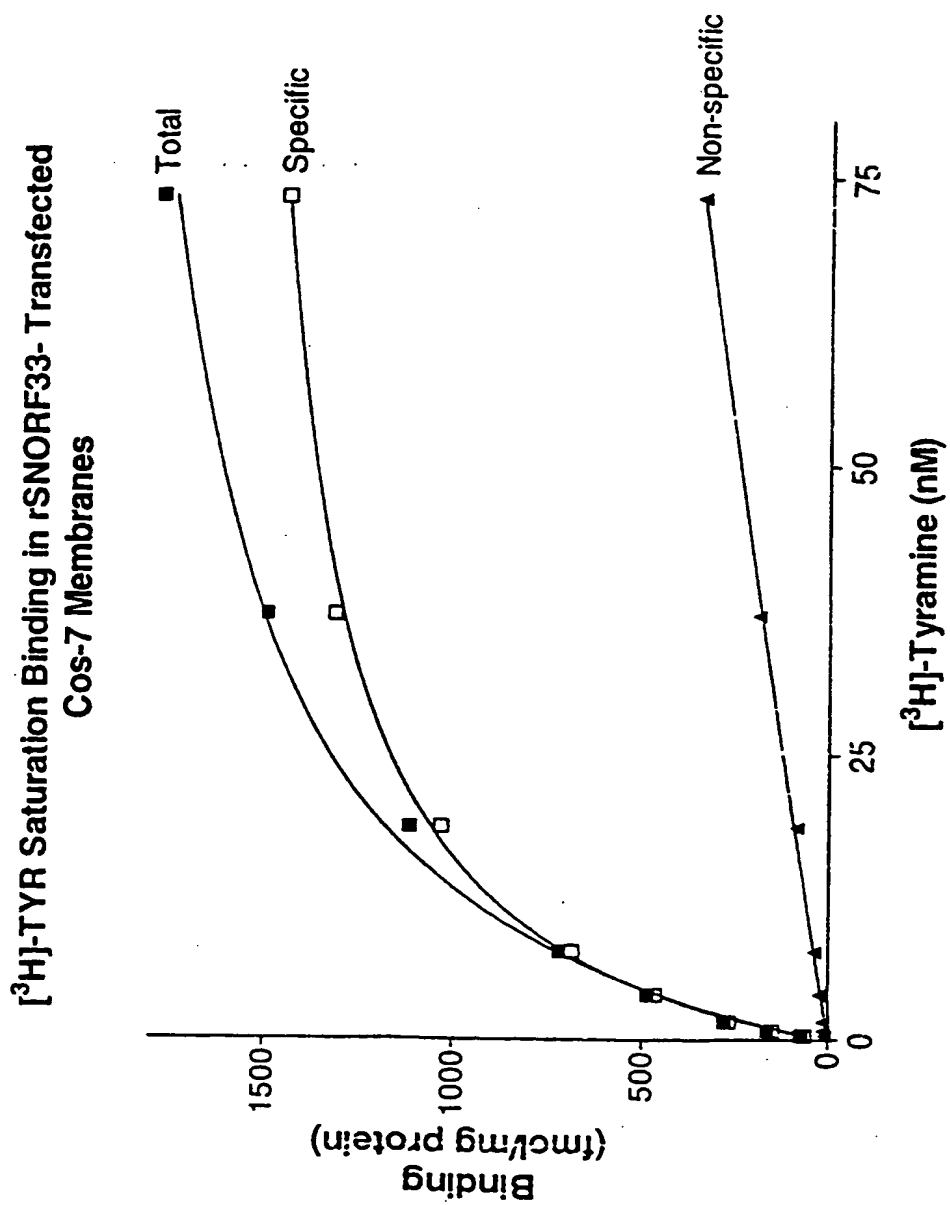






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Figure 13



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Figure 14

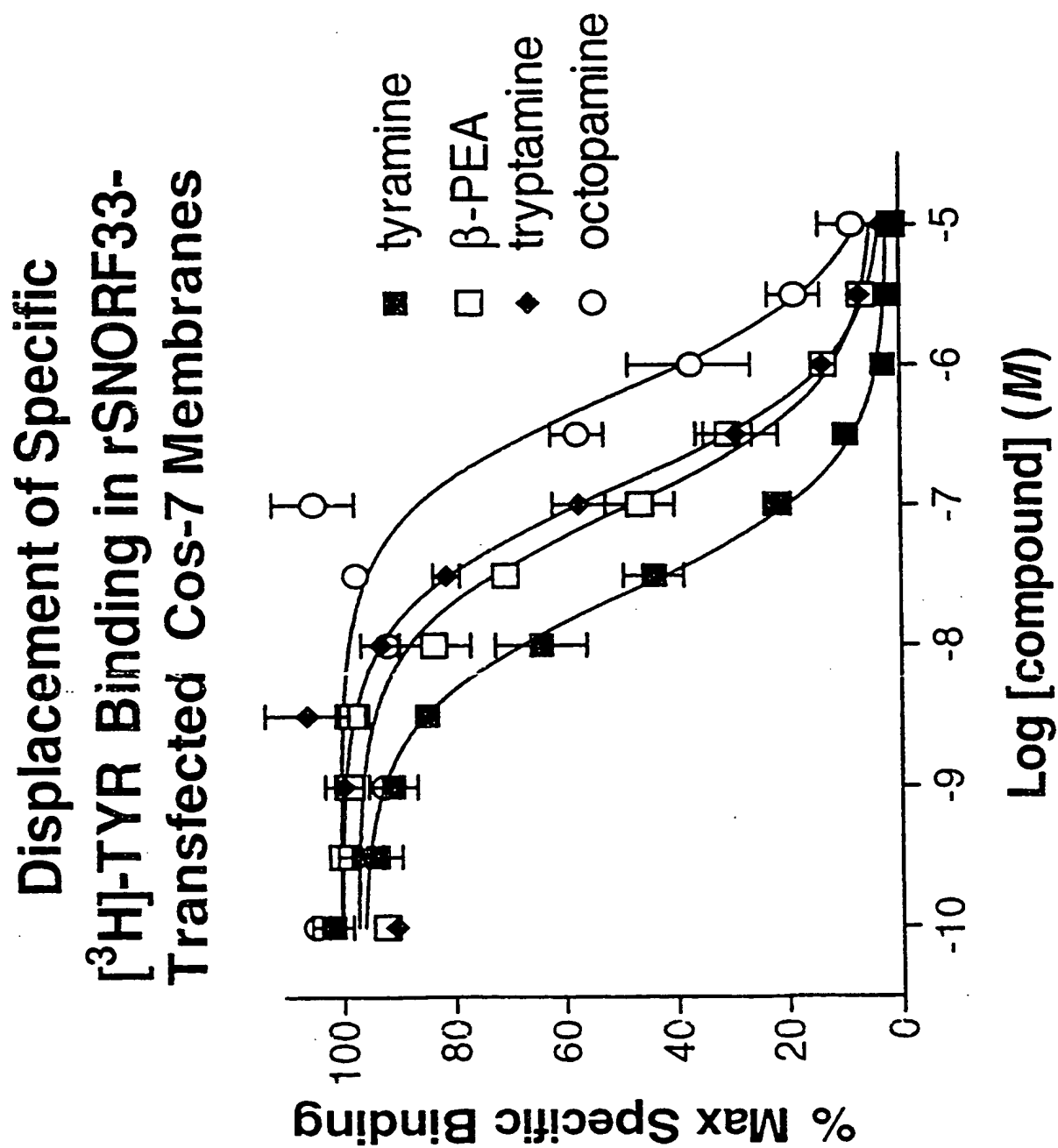
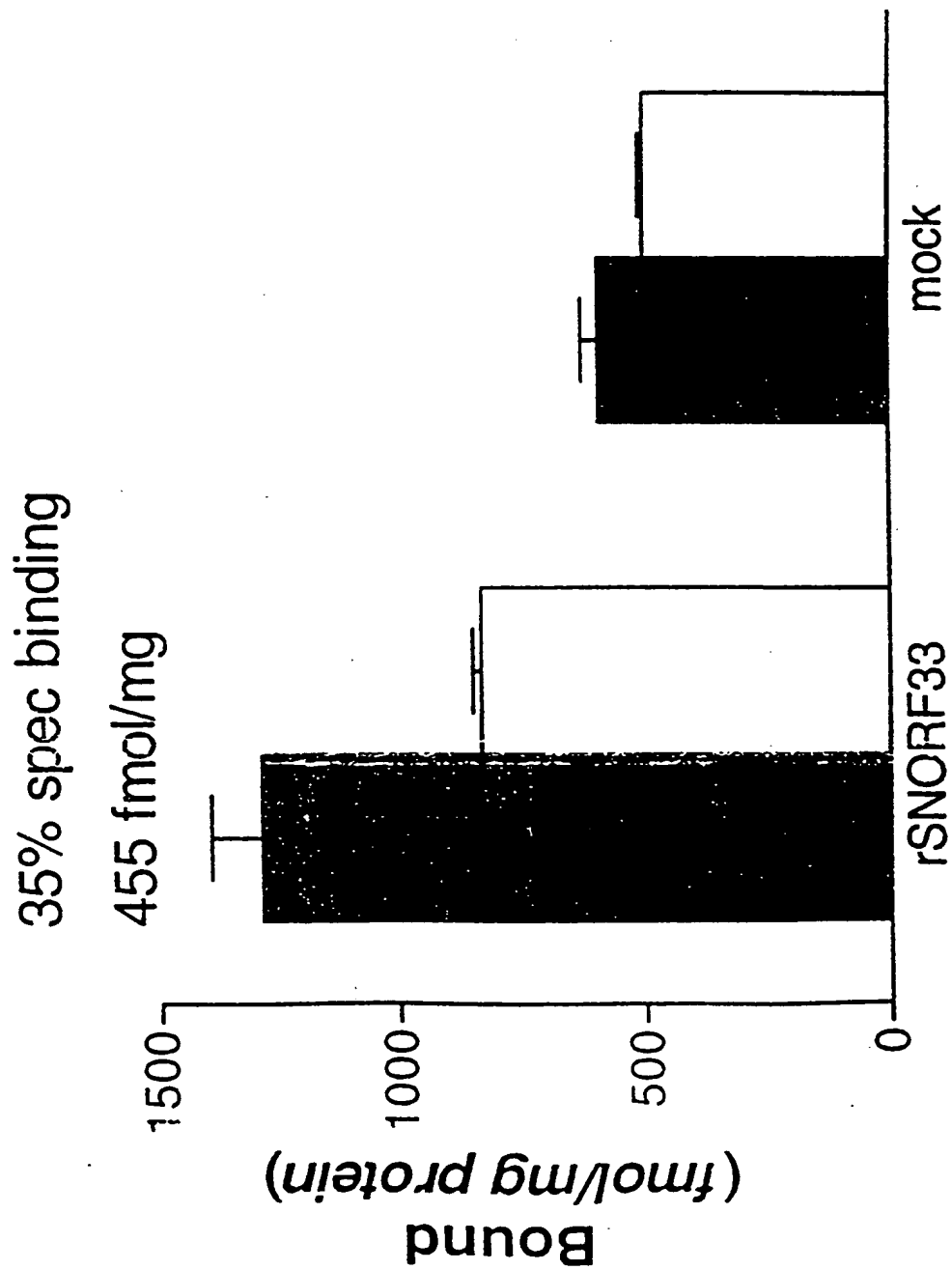


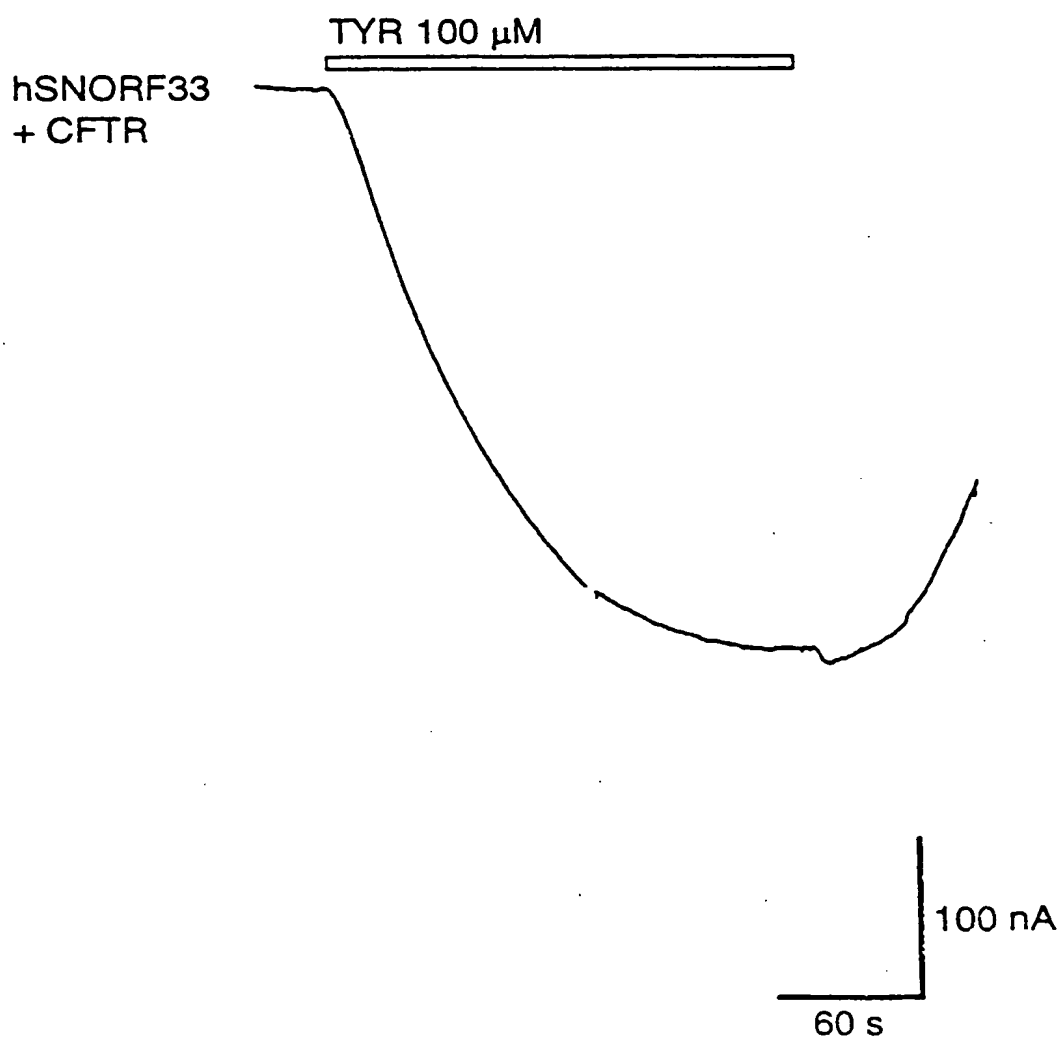
Figure 15

# [<sup>3</sup>H]-T (20 nM) Binding in rSNORF33- and Mock- Transfected Cos-7 Membranes



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Figure 16



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Figure 17

|     |   |     |
|-----|---|-----|
| 1   | GGTACTGGCGTTTCATGACTTCCCTTCTATATATACCTGGATCTGTATATGTTATTGTTTACTTA | 60  |
| 61  | TAGGATATATTTTCATAGCTTAAAGGACAAAGCAAGGTCAATCAATCGTACGAAATGTTTCAAGT | 120 |
| 121 | TGGATTGGAAGGAAAGCCCAAGCACCAAGCAAGGAACAAGCCGCCGAAGACCTT            | 180 |
| 181 | AGGATCATGGTGGCGGTTTTCCTTCGTATGCTGGTGGCCCGTTCTTCTCTGCACGGTCCT      | 240 |
| 241 | GGACCCCTTTCCT   | 252 |

## Figure 18

81 D P F  
83

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## FIGURE 19A

1 TGCAGTGATGATCCTTTTGCCACGCTATCACAAACATTTCCACAGAAACAGCGACTGGTC 60

61 AAGAGAAAGTCCAAGCTTCCCTGTACAGCTTAATGTCACTCATAATCCTGGCCACTCTGGT 120

121 TGGCAACTTAATAGTAATAATTCCATATCCCATTTCAAGCAACTTCATACACCCACCAA 180

181 CTGGCTCCTTCACCTCCATGGCCATTGTGCGACTTTCTGCTGGGCTGTCTGATAATGCCCTG 240

241 CAGCATGGTGAGAACTGTTGAGCGCTGTGGTATTTTGGGAAATCCTCTGTAAAGTTCA 300

301 CACCAGCACCGATATCATGCTGAGCTCCGCCCTCCATTTTCCACTTAGCTTTCATTTCCAT 360

361 TGACCGCTACTGTGCTGTGACCCCTTTGAGATACAAAGCCAAGATCAATATCTCCAC 420

421 TATTCTTGATGATCCCTCGTTAGTTGGAGCCTTCCCTGCTGTTTATGCATTTGGGATGAT 480

481 CTTCCTGGAACTGAACCTTAAAGGAGTGGAAGAGCTGTATCGCAGTCAGGTCAGCGACCT 540

541 GGGCGGCTGTTCTCCCTTCTTTAGTAAAGTATCTGGGGTACTGGCGTTCATGACTTCCCTT 600



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## FIGURE 19B

601 CTATATACCTGGATCTGTTATGTTATTTGTTTACTATAGGATATATTTTCATAGCTAAAGG 660

661 ACAAGCAAGTCAATCAATCGTACGAATGTTCAAGTTGGATTGGAAGGAAAGCCAAGC 720

721 ACCACAAAGCAAGAAACAAAGCCGGAAGACCTTAGGGATCATGGTGGCGGTTTTCCT 780

781 CGTATGCTGGTGCCCGTTCTTCTCTGCACGGTCCTGGACCCCTTTCCTGGGCTATGTTAT 840

841 CCCACCCCTCTCTGAATGACGCACTGTATTGGTTTGGGTACTTGAATTCTGCCCCCAATCC 900

901 GATGGTTTATGCCCTTTTCTATATCCCTGGTTCAGAAAGAGCCCTTGAAGATGGTTCTCCTTGG 960

961 TAAAAATTTCCAAAAAGATTCACTAGGTCTAAGCTATTTTGTAAACGCAATTCATGAAA 1020

1021 CCCATGTATTT 1031

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FIGURE 20A

|     |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |     |
|-----|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|-----|
| 1   | M | H | L | C | H | A | I | T | N | I | S | H | R | N | S | D | W | S | R | E | 20  |
| 21  | V | Q | A | S | L | Y | S | L | M | S | L | I | I | L | A | T | L | V | G | N | 40  |
| 41  | L | I | V | I | I | S | I | S | H | F | K | Q | L | H | T | P | T | N | W | L | 60  |
| 61  | L | H | S | M | A | I | V | D | F | L | L | G | C | L | I | M | P | C | S | M | 80  |
| 81  | V | R | T | V | E | R | C | W | Y | F | G | E | I | L | C | K | V | H | T | S | 100 |
| 101 | T | D | I | M | L | S | S | A | S | I | F | H | L | A | F | I | S | I | D | R | 120 |
| 121 | Y | C | A | V | C | D | P | L | R | Y | K | A | K | I | N | I | S | T | I | L | 140 |
| 141 | V | M | I | L | V | S | W | S | L | P | A | V | Y | A | F | G | M | I | F | L | 160 |
| 161 | E | L | N | L | K | G | V | E | E | L | Y | R | S | Q | V | S | D | L | G | G | 180 |



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## FIGURE 21

|           |         |            |            |            |            |
|-----------|---------|------------|------------|------------|------------|
| Rat       | SNORF33 | -MhlCHnsaN | IShtnsnWSr | dVrASLYSLi | sLIILtTLVG |
| Mouse     | SNORF33 | -MhlCHaitN | IShrnsdWSr | eVqASLYSLm | sLIILaTLVG |
| Human     | SNORF33 | mMpfCHniiN | IScvknnWSn | dVrASLYSLm | vLIILtTLVG |
| Consensus |         | -M--CH---N | IS-----WS- | -V-ASLYSL- | -LIIL-TLVG |

|           |         |            |            |            |            |
|-----------|---------|------------|------------|------------|------------|
| Rat       | SNORF33 | NLIViisISH | FKQLHTPTNW | LlHSMAvVDF | LLGCLvMPys |
| Mouse     | SNORF33 | NLIViisISH | FKQLHTPTNW | LlHSMAiVDF | LLGCLiMPcs |
| Human     | SNORF33 | NLIVivSISH | FKQLHTPTNW | LiHSMAvVDF | LLGCLvMPys |
| Consensus |         | NLIVI-SISH | FKQLHTPTNW | L-HSMA-VDF | LLGCL-MP-S |

|           |         |            |            |            |            |
|-----------|---------|------------|------------|------------|------------|
| Rat       | SNORF33 | MVRtvEhCWY | FGElfCKlHT | STDIMLSSAS | IlHLaFISID |
| Mouse     | SNORF33 | MVRtvErCWY | FGEilCKvHT | STDIMLSSAS | IfHLaFISID |
| Human     | SNORF33 | MVRsaEhCWY | FGEvfCKiHT | STDIMLSSAS | IfHLsFISID |
| Consensus |         | MVR--E-CWY | FGE--CK-HT | STDIMLSSAS | I-HL-FISID |

|           |         |            |            |            |            |
|-----------|---------|------------|------------|------------|------------|
| Rat       | SNORF33 | RYyAVCDPLR | YKAKiNlaaI | fVMiliSWSl | PAVfAFGMIF |
| Mouse     | SNORF33 | RYcAVCDPLR | YKAKiNistI | lVMilvSWSl | PAVyAFGMIF |
| Human     | SNORF33 | RYyAVCDPLR | YKAKmNilvI | cVMIfiSWSv | PAVfAFGMIF |
| Consensus |         | RY-AVCDPLR | YKAK-N---I | -VMI--SWS- | PAV-AFGMIF |

|           |         |            |            |            |            |
|-----------|---------|------------|------------|------------|------------|
| Rat       | SNORF33 | LELNleGvEE | lYhnqVfclr | GCfpFFSKvS | GVLafMTSFY |
| Mouse     | SNORF33 | LELNlkGvEE | lYrsqVsdlg | GCspFFSKvS | GVLafMTSFY |
| Human     | SNORF33 | LELNfkGaEE | iYykhVhcrG | GCsvFFSKiS | GVLtFMTSFY |
| Consensus |         | LELN--G-EE | -Y---V---- | GC--FFSK-S | GVL-FMTSFY |

|           |         |            |            |            |            |
|-----------|---------|------------|------------|------------|------------|
| Rat       | SNORF33 | IPGSvMLfVY | YRIYfiAKgQ | ARsInraN.. | lQvGLEgesr |
| Mouse     | SNORF33 | IPGSvMLfVY | YRIYfiAKgQ | ARsInrtN.. | vQvGLEgksq |
| Human     | SNORF33 | IPGSiMLcVY | YRIYliAKeQ | ARlIsdaNqk | lQiGLEmknG |
| Consensus |         | IPGS-ML-VY | YRIY-IAK-Q | AR-I---N-- | -Q-GLE---- |

|           |         |            |            |            |            |
|-----------|---------|------------|------------|------------|------------|
| Rat       | SNORF33 | apQSKEtKAa | KTLGImvGVF | LlCWCPFFfC | mVlDPFLgYv |
| Mouse     | SNORF33 | apQSKEtKAa | KTLGImvGVF | LvCWCPFFlC | tVlDPFLgYv |
| Human     | SNORF33 | isQSKErKAv | KTLGIvmGVF | LiCWCPFFiC | tVmDPFLhYi |
| Consensus |         | --QSKE-KA- | KTLGI--GVF | L-CWCPFF-C | -V-DPFL-Y- |

|           |         |            |             |            |            |
|-----------|---------|------------|-------------|------------|------------|
| Rat       | SNORF33 | IPPtLNDtLn | WFGYLNSafN  | PMVYAFFYPW | FRrALKMvLf |
| Mouse     | SNORF33 | IPPsLNDaLy | WFGYLNSalN  | PMVYAFFYPW | FRrALKMvLl |
| Human     | SNORF33 | IPPtLNDvLi | WFGYLNSstfn | PMVYAFFYPW | FRkALKMmLf |
| Consensus |         | IPP-LND-L- | WFGYLNS--N  | PMVYAFFYPW | FR-ALKM-L- |

|           |         |            |           |
|-----------|---------|------------|-----------|
| Rat       | SNORF33 | GKIFQKDSSR | sKLFL     |
| Mouse     | SNORF33 | GKIFQKDSSR | sKLFL     |
| Human     | SNORF33 | GKIFQKDSSR | cKLFLelss |
| Consensus |         | GKIFQKDSSR | -KLFL---- |